

Appl. No. : 10/007,768
Filed : November 8, 2001

motif (whose amino acid sequence is as follows: SLKLMATLFSTYASA (SEQ ID NO:1)). This motif is found in proteins from a multitude of viruses, bacteria, fungi, and protozoa, which are involved in cardiovascular disease.

Please replace Paragraph [0055] as follows:

[0055] The following peptides were synthesized by a robotic multiple peptide synthesizer:

Infectious agent	Peptide portion
1. Chlamydia pneumoniae	LPTAVLNLTAWNPSLLGNATALST (SEQ ID NO:2)
2. Mycoplasma pneumoniae	TPPNMATLVSTAMSL (SEQ ID NO:3)
3. Borrelia burgdorferi	LFLIMATFLSPSISG (SEQ ID NO:4)
4. Treponema pallidum	RSEAMALVLSTLENR (SEQ ID NO:5)
5. Trypanozoma cruzi	NTFHMAGGGSTLINL (SEQ ID NO:6)
6. Cocksackie virus	FIEWLKVKILPEVKEKHEFLSRL (SEQ ID NO:7)
7. Epstein-Barr virus	TPSPAIPSHSSNTALERPLA (SEQ ID NO:8)
8. Cytomegalovirus	VMAPRTLILTVGLLCMRI (SEQ ID NO:9)
9. Helicobacter pylori	TYNBMATGTSPVMSG (SEQ ID NO:10)
10. Streptococcus Group-A	RVTTRSQAQDAAGLKEKADC (SEQ ID NO:11)

IN THE SEQUENCE LISTING:

Please add the attached Sequence Listing pages 1-3.

REMARKS

This Preliminary Amendment brings the patent application into compliance with the Sequence Listing Disclosure requirement of the United States Patent and Trademark Office. Enclosed herewith are: (1) a paper copy of the Sequence Listing, (2) and a computer readable version of the Sequence Listing. The Preliminary Amendment directs entry of the paper copy of the Sequence Listing in to the application. In view of the foregoing, the application is believed to fully comply with the Sequence Listing Disclosure requirements.

The specification has been amended to include the sequence identification numbers from the sequence listing. Accordingly, no new matter has been added. The specific changes to the